

SEQUENCE LISTING

<110> Friddle, Carl Johan
Aylor, Erin
Scoville, John
Walke, D. Wade

<120> Novel Human Secreted Signal Proteins and Polynucleotides Encoding the Same

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<150> US 60/216,384
<151> 2000-07-07

<150> US 60/219,890
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<213> homo sapiens

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ttcccgcc	gccggtgaa	ctgcaccacc	gtccacgaca	gcctggccat	cttcgggccc	300
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Ser Ser Leu Gly Ser Gln Pro Ile Leu Cys Ala Ser Ile Pro Gly Leu
35 40 45
Val Pro Lys Gln Leu Arg Phe Cys Arg Asn Tyr Val Glu Ile Met Pro
50 55 60
Ser Val Ala Glu Gly Ile Lys Ile Gly Ile Gln Glu Cys Gln His Gln
65 70 75 80
Phe Arg Gly Arg Arg Trp Asn Cys Thr Thr Val His Asp Ser Leu Ala
85 90 95
Ile Phe Gly Pro Val Leu Asp Lys Ala Thr Arg Glu Ser Ala Phe Val
100 105 110
His Ala Ile Ala Ser Ala Gly Val Ala Phe Ala Val Thr Arg Ser Cys
115 120 125
Ala Glu Gly Thr Ala Ala Ile Cys Gly Cys Ser Ser Arg His Gln Gly
130 135 140
Ser Pro Gly Lys Gly Trp Lys Trp Gly Gly Cys Ser Glu Asp Ile Glu
145 150 155 160
Phe Gly Gly Met Val Ser Arg Glu Phe Ala Asp Ala Arg Glu Asn Arg
165 170 175
Pro Asp Ala Arg Ser Ala Met Asn Arg His Asn Asn Glu Ala Gly Arg
180 185 190
Gln Ala Ile Ala Ser His Met His Leu Lys Cys Lys Cys His Gly Leu
195 200 205
Ser Gly Ser Cys Glu Val Lys Thr Cys Trp Trp Ser Gln Pro Asp Phe
210 215 220
Arg Ala Ile Gly Asp Phe Leu Lys Asp Lys Tyr Asp Ser Ala Ser Glu
225 230 235 240
Met Val Val Glu Lys His Arg Glu Ser Arg Gly Trp Val Glu Thr Leu
245 250 255
Arg Pro Arg Tyr Thr Tyr Phe Lys Val Pro Thr Glu Arg Asp Leu Val
260 265 270
Tyr Tyr Glu Ala Ser Pro Asn Phe Cys Glu Pro Asn Pro Glu Thr Gly
275 280 285
Ser Phe Gly Thr Arg Asp Arg Thr Cys Asn Val Ser Ser His Gly Ile
290 295 300
Asp Gly Cys Asp Leu Leu Cys Cys Gly Arg Gly His Asn Ala Arg Ala
305 310 315 320
Glu Arg Arg Arg Glu Lys Cys Arg Cys Val Phe His Trp Cys Cys Tyr
325 330 335
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340 345 350
Asp Gly Cys Leu Arg Thr Gly His Ser Gly Pro Cys Arg Ser Leu Ala
355 360 365
Trp Ile Trp Ser Pro Gly Ser Gln Gly His Asp Leu Leu Glu Gln Leu
370 375 380
Pro Arg Ser Gly Gly Leu Gly Gln Cys Ser Ser Leu Gln Asn Trp Thr
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 Ser Ser Leu Gly Ser Gln Pro Ile Leu Cys Ala Ser Ile Pro Gly Leu
 35 40 45
 Val Pro Lys Gln Leu Arg Phe Cys Arg Asn Tyr Val Glu Ile Met Pro
 50 55 60
 Ser Val Ala Glu Gly Ile Lys Ile Gly Ile Gln Glu Cys Gln His Gln
 65 70 75 80
 Phe Arg Gly Arg Arg Trp Asn Cys Thr Thr Val His Asp Ser Leu Ala
 85 90 95
 Ile Phe Gly Pro Val Leu Asp Lys Ala Thr Arg Glu Ser Ala Phe Val
 100 105 110
 His Ala Ile Ala Ser Ala Gly Val Ala Phe Ala Val Thr Arg Ser Cys
 115 120 125
 Ala Glu Gly Thr Ala Ala Ile Cys Gly Cys Ser Ser Arg His Gln Gly
 130 135 140
 Ser Pro Gly Lys Gly Trp Lys Trp Gly Gly Cys Ser Glu Asp Ile Glu
 145 150 155 160
 Phe Gly Gly Met Val Ser Arg Glu Phe Ala Asp Ala Arg Glu Asn Arg

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Ser Gly Ser Cys Glu Val Lys Thr Cys Trp Trp Ser Gln Pro Asp Phe			
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Arg Ala Ile Gly Asp Phe Leu Lys Asp Lys Tyr Asp Ser Ala Ser Glu			
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Met Val Val Glu Lys His Arg Glu Ser Arg Gly Trp Val Glu Thr Leu			
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260	265	270	
Tyr Tyr Glu Ala Ser Pro Asn Phe Cys Glu Pro Asn Pro Glu Thr Gly			
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Ser Phe Gly Thr Arg Asp Arg Thr Cys Asn Val Ser Ser His Gly Ile			
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Asp Gly Cys Asp Leu Leu Cys Cys Gly Arg Gly His Asn Ala Arg Ala			
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<213> homo sapiens

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															20
															25
															30
Cys	Leu	Thr	Phe	Ser	Leu	Phe	Gly	Arg	Ser	Val	Asn	Asn	Phe	Leu	Ile
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															40
															45
Thr	Gly	Pro	Lys	Ala	Tyr	Leu	Thr	Tyr	Thr	Thr	Ser	Val	Ala	Leu	Gly
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															55
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Ala	Gln	Ser	Gly	Ile	Glu	Glu	Cys	Lys	Phe	Gln	Phe	Ala	Trp	Glu	Arg
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Trp	Asn	Cys	Pro	Glu	Asn	Ala	Leu	Gln	Leu	Ser	Thr	His	Asn	Arg	Leu
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Arg	Ser	Ala	Thr	Arg	Glu	Thr	Ser	Phe	Ile	His	Ala	Ile	Ser	Ser	Ala
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Gly	Val	Met	Tyr	Ile	Ile	Thr	Lys	Asn	Cys	Ser	Met	Gly	Asp	Phe	Glu
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Leu	Met	Asn	Leu	His	Asn	Asn	Arg	Ala	Gly	Arg	Leu	Ala	Val	Arg	Ala
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 Ile Gln Thr Cys Trp Leu Gln Leu Ala Glu Phe Arg Glu Met Gly Asp
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 225 230 235 240
 Arg Gln Leu Arg Ala Gly Asn Ser Ala Glu Gly His Trp Val Pro Ala
 245 250 255
 Glu Ala Phe Leu Pro Ser Ala Glu Ala Glu Leu Ile Phe Leu Glu Glu
 260 265 270
 Ser Pro Asp Tyr Cys Thr Cys Asn Ser Ser Leu Gly Ile Tyr Gly Thr
 275 280 285
 Glu Gly Arg Glu Cys Leu Gln Asn Ser His Asn Thr Ser Arg Trp Glu
 290 295 300
 Arg Arg Ser Cys Gly Arg Leu Cys Thr Glu Cys Gly Leu Gln Val Glu
 305 310 315 320
 Glu Arg Lys Thr Glu Val Ile Ser Ser Cys Asn Cys Lys Phe Gln Trp
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 Ala

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gaacgcgtga	actgcccctga	aaatgctctt	cagctctcca	cccacaacag	gctgagaagt	240
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accaagaact	gtagcatggg	tgacttcgaa	aactgtggct	gtgatgggtc	aaacaatgga	360
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Pro	Lys	Ala	Tyr	Leu	Thr	Tyr	Thr	Thr	Ser	Val	Ala	Leu	Gly	Ala	Gln
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Ser	Gly	Ile	Glu	Glu	Cys	Lys	Phe	Gln	Phe	Ala	Trp	Glu	Arg	Trp	Asn
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Cys	Pro	Glu	Asn	Ala	Leu	Gln	Leu	Ser	Thr	His	Asn	Arg	Leu	Arg	Ser
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Met	Tyr	Ile	Ile	Thr	Lys	Asn	Cys	Ser	Met	Gly	Asp	Phe	Glu	Asn	Cys
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Gly	Cys	Asp	Gly	Ser	Asn	Asn	Gly	Lys	Thr	Gly	Gly	His	Gly	Trp	Ile
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Trp	Gly	Gly	Cys	Ser	Asp	Asn	Val	Glu	Phe	Gly	Glu	Arg	Ile	Ser	Lys
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Leu	Phe	Val	Asp	Ser	Leu	Glu	Lys	Gly	Lys	Asp	Ala	Arg	Ala	Leu	Met
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Asn	Leu	His	Asn	Asn	Arg	Ala	Gly	Arg	Leu	Ala	Val	Arg	Ala	Thr	Met
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Lys	Arg	Thr	Cys	Lys	Cys	His	Gly	Ile	Ser	Gly	Ser	Cys	Ser	Ile	Gln
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Thr	Cys	Trp	Leu	Gln	Leu	Ala	Glu	Phe	Arg	Glu	Met	Gly	Asp	Tyr	Leu
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Lys	Ala	Lys	Tyr	Asp	Gln	Ala	Leu	Lys	Ile	Glu	Met	Asp	Lys	Arg	Gln
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Leu	Arg	Ala	Gly	Asn	Ser	Ala	Glu	Gly	His	Trp	Val	Pro	Ala	Glu	Ala
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Phe	Leu	Pro	Ser	Ala	Glu	Ala	Glu	Leu	Ile	Phe	Leu	Glu	Glu	Ser	Pro
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Asp	Tyr	Cys	Thr	Cys	Asn	Ser	Ser	Leu	Gly	Ile	Tyr	Gly	Thr	Glu	Gly
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Arg	Glu	Cys	Leu	Gln	Asn	Ser	His	Asn	Thr	Ser	Arg	Trp	Glu	Arg	Arg
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Ser	Cys	Gly	Arg	Leu	Cys	Thr	Glu	Cys	Gly	Leu	Gln	Val	Glu	Glu	Arg
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Lys	Thr	Glu	Val	Ile	Ser	Ser	Cys	Asn	Cys	Lys	Phe	Gln	Trp	Cys	Cys
					305			310						320	
Thr	Val	Lys	Cys	Asp	Gln	Cys	Arg	His	Val	Val	Ser	Lys	Tyr	Tyr	Cys
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<213> homo sapiens

<400> 11

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Leu Phe Val Asp Ser Leu Glu Lys Gly Lys Asp Ala Arg Ala Leu Met	
50 55 60	
Asn Leu His Asn Asn Arg Ala Gly Arg Leu Ala Val Arg Ala Thr Met	
65 70 75 80	
Lys Arg Thr Cys Lys Cys His Gly Ile Ser Gly Ser Cys Ser Ile Gln	
85 90 95	
Thr Cys Trp Leu Gln Leu Ala Glu Phe Arg Glu Met Gly Asp Tyr Leu	
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130 135 140	
Phe Leu Pro Ser Ala Glu Ala Glu Leu Ile Phe Leu Glu Glu Ser Pro	
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165 170 175	
Arg Glu Cys Leu Gln Asn Ser His Asn Thr Ser Arg Trp Glu Arg Arg	
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Ser Cys Gly Arg Leu Cys Thr Glu Cys Gly Leu Gln Val Glu Glu Arg	
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Lys Thr Glu Val Ile Ser Ser Cys Asn Cys Lys Phe Gln Trp Cys Cys	
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<213> homo sapiens

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<211> 34
<212> PRT
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Pro Ile

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<212> DNA
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<210> 17
<211> 36
<212> PRT
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<212> DNA

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<211> 351

<212> PRT

<213> homo sapiens

<400> 19

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Pro Lys Ala Tyr Leu Thr Tyr Thr Ser Val Ala Leu Gly Ala Gln
35 40 45
Ser Gly Ile Glu Glu Cys Lys Phe Gln Phe Ala Trp Glu Arg Trp Asn
50 55 60
Cys Pro Glu Asn Ala Leu Gln Leu Ser Thr His Asn Arg Leu Arg Ser
65 70 75 80
Ala Thr Arg Glu Thr Ser Phe Ile His Ala Ile Ser Ser Ala Gly Val
85 90 95
Met Tyr Ile Ile Thr Lys Asn Cys Ser Met Gly Asp Phe Glu Asn Cys
100 105 110
Gly Cys Asp Gly Ser Asn Asn Gly Lys Thr Gly Gly His Gly Trp Ile
115 120 125
Trp Gly Gly Cys Ser Asp Asn Val Glu Phe Gly Glu Arg Ile Ser Lys
130 135 140
Leu Phe Val Asp Ser Leu Glu Lys Gly Lys Asp Ala Arg Ala Leu Met
145 150 155 160
Asn Leu His Asn Asn Arg Ala Gly Arg Leu Ala Val Arg Ala Thr Met
165 170 175
Lys Arg Thr Cys Lys Cys His Gly Ile Ser Gly Ser Cys Ser Ile Gln
180 185 190
Thr Cys Trp Leu Gln Leu Ala Glu Phe Arg Glu Met Gly Asp Tyr Leu
195 200 205

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Lys Ala Lys Tyr Asp Gln Ala Leu Lys Ile Glu Met Asp Lys Arg Gln
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 Leu Arg Ala Gly Asn Ser Ala Glu Gly His Trp Val Pro Ala Glu Ala
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 Phe Leu Pro Ser Ala Glu Ala Glu Leu Ile Phe Leu Glu Glu Ser Pro
 245 250 255
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 260 265 270
 Arg Glu Cys Leu Gln Asn Ser His Asn Thr Ser Arg Trp Glu Arg Arg
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 Ser Cys Gly Arg Leu Cys Thr Glu Cys Gly Leu Gln Val Glu Glu Arg
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 Lys Thr Glu Val Ile Ser Ser Cys Asn Cys Lys Phe Gln Trp Cys Cys
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<210> 22
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 <212> DNA
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<210> 23
 <211> 36
 <212> PRT
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<400> 23

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 20 25 30
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<212> DNA

<213> homo sapiens

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